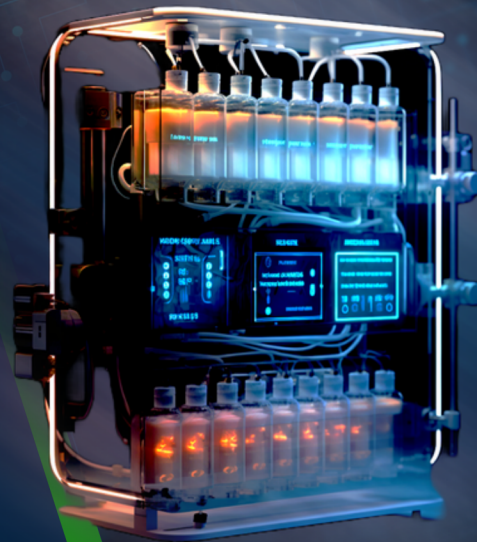




# OPTIMIZATION

The Optimization Module performs the multidimensional search needed to fit model parameters to one or more data sets automatically. Objective function weighting is user-defined, and includes the most common weighting schemes.



## Fitting Models to Data

One of the most important uses of GastroPlus® is to fit absorption, pharmacokinetic, and pharmacodynamic models to observations. In doing so, researchers gain tremendous insight into how their compound is behaving *in vivo*. When a single set of model parameters can be found that properly describes the observed plasma concentration vs. time profiles for all dose levels, a useful model has been obtained. In general, if the model parameters must be changed for each dose level, then something is not being accounted for correctly.

## Example Parameters:

- ✓ PBPK model parameters to plasma and/or tissue concentration vs. time data
- ✓  $P_{eff}$  and absorption scale factors to determine regional dependencies
- ✓ A wide variety of physiological parameters (when necessary – rarely used)
- ✓ Parameters to match profiles of parent drugs or any of their metabolites

## What is the Optimization module?

The Optimization Module for GastroPlus extends and enhances the program's basic capabilities in several important ways:

- ✓ To automatically fit physiochemical, pharmacokinetic, formulation and/or physiology model parameters to data
- ✓ To optimize study designs (e.g., dosing regimens)
- ✓ To "deconvolute" *in vivo* release profiles to achieve a target plasma (or tissue) concentration vs. time curves for parent and/or metabolites



### Flexible optimization settings

Optimization with multiple drug records, multiple objective functions, and multiple observed data like concentration profile,  $C_{max}$ ,  $T_{max}$ , and AUC.

